SEQUENCE LISTING

<110> Wei, Ying-Fei et al.

<120> Chemokine Alpha-6

<130> PF458D1

<150> 09/177,304
<151> 1998-10-23

<150> 60/063,387
<151> 1997-10-24

<150> 60/079,245
<151> 1998-03-25

<160> 18

<170> PatentIn Ver. 3.1

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<212> DNA
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<222> (46)..(297)

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Phe Leu Ser Val Pro Cys Leu Leu Leu Leu Pro Ala Arg Val Val Trp
  5                  10                  15                  20

ggg tgt tgg tgt ttc ctc cct gga gaa gat ggg gga ggc tgt ccc act   153
Gly Cys Trp Cys Phe Leu Pro Gly Glu Asp Gly Gly Gly Cys Pro Thr
                 25                  30                  35

ccc agc tct ggc aga atc aag ctg ttg cag cag tgc ctt ctt cat cct   201
Pro Ser Ser Gly Arg Ile Lys Leu Leu Gln Gln Cys Leu Leu His Pro
             40                  45                  50

tcc tta cga tca atc aca gtc tcc aga aga tca gct caa ttg ctg tgc   249
Ser Leu Arg Ser Ile Thr Val Ser Arg Arg Ser Ala Gln Leu Leu Cys
         55                  60                  65

agg tta aaa cta cag aac cac atc cca aag gta cct ggt aag aat gtt   297
Arg Leu Lys Leu Gln Asn His Ile Pro Lys Val Pro Gly Lys Asn Val
     70                  75                  80

tgaaagatct tccatttcta ggaaccccag tcctgcttct ccgcaatggc acatgcttcc 357

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Arg Val Val Trp Gly Cys Trp Cys Phe Leu Pro Gly Glu Asp Gly Gly
20 25 30

Gly Cys Pro Thr Pro Ser Ser Gly Arg Ile Lys Leu Leu Gln Gln Cys
35 40 45

Leu Leu His Pro Ser Leu Arg Ser Ile Thr Val Ser Arg Arg Ser Ala
50 55 60

Gln Leu Leu Cys Arg Leu Lys Leu Gln Asn His Ile Pro Lys Val Pro
65 70 75 80

Gly Lys Asn Val

<210> 3
<211> 75
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<400> 3
Cys Lys Cys Ser Arg Lys Gly Pro Lys Ile Arg Tyr Ser Asp Val Lys
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Lys Leu Glu Met Lys Pro Lys Tyr Pro His Cys Glu Glu Lys Met Val
20 25 30

Ile Ile Thr Thr Lys Ser Val Ser Arg Tyr Arg Gly Gln Glu His Cys
35 40 45

Leu His Pro Lys Leu Gln Ser Thr Lys Arg Phe Ile Lys Trp Tyr Asn
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Val Ala Ser Leu Lys Asn Gly Lys Gln Val Cys Leu Asp Pro Glu Ala
35 40 45

Pro Phe Leu Lys Lys Val Ile Gln Lys Ile Leu Asp Ser Gly Thr Arg
50 55 60

Asn
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20 25 30

Val Ile Ile Thr Leu Lys Glu Asn Lys Gly Gln Arg Cys Leu Asn Pro
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cagaaccaca tcccaaaggt acctggtaag aatgtttgaa agatcttcca tttctaggaa 180

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cagaaccaca tcccaaaggt acctggtaag aatgtttgaa agatcttcca tttctaggaa 180

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gtacctttgg gatgtggttc tgtagtttta acctgcacag caattgagct gatcttctgg 180

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gtacctttgg gatgtggttc tgtagtttta acctgcacag caattgagct gatcttctgg 180

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<223> n equals a, t, g, or c

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cccagtcctg cttctccgca atggcacatg cttccactcc atccatactg gcatcctcaa 120

ataaacagat atgtatacat at 142

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atccatactg gcatcctcaa ataaacagat atgtatacat ataaa 105

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tacctttggg atgtggttct gtagttttaa cctgcacagc aattgagctg atcttctgga 180

gactgtgatt gatcgtaagg aaggatggag aaggcactgc tgcaacagct tgattctgcc 240

agagctggga gtgggacagc ctcccccatc ttctccaggg aggaaacacc aacacccca 300

aaccacccgg gcaggtaaga ggagcaagca cggcacagag aggaagggcc tctgcatttt 360

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ctccctggna gaagatgggg gaggctgtcc cactcccagc tctggtcagg aatgcaagnt 180

gttggcagca gtgnccttct tgcatgcctt gccttnacgg atgcaatgca cagtgctccc 240

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caacatgccc aaaggttaac ctgggttcaa agaaatgttt ttgna 345

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<210> 16
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<223> Contains a BamHI restriction enzyme site and an efficient signal for initiation of translation in eukaryotic cells

<400> 17
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